

SEQUENCE LISTING

<110> Pan, Yang

<120> NOVEL MOLECULES OF THE TANGO-93-RELATED
PROTEIN FAMILY AND USES THEREOF

<130> 07334-369001

<140> US 10/134,410

<141> 2002-04-29

<150> US 09/131,263

<151> 1998-08-07

<150> US 09/369,693

<151> 1999-08-06

<160> 14

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 1360

<212> DNA

<213> Mus musculus

<220>

<221> CDS

<222> (137)...(604)

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acatactgtg gagctc atg atg gtt ctg agt ggg gca cta tgc ttc cga atg	172
Met Met Val Leu Ser Gly Ala Leu Cys Phe Arg Met	
1 5 10	

aag gat tca gcc ttg aag gta ctg tat ctg cac aat aac cag ctg ctg	220
Lys Asp Ser Ala Leu Lys Val Leu Tyr Leu His Asn Asn Gln Leu Leu	
15 20 25	

gct gga gga ctg cac gca gag aag gtc att aaa ggt gag gag atc agt	268
Ala Gly Gly Leu His Ala Glu Lys Val Ile Lys Gly Glu Glu Ile Ser	
30 35 40	

gtt gtc cca aat cgg gca ctg gat gcc agt ctg tcc cct gtc atc ctg	316
Val Val Pro Asn Arg Ala Leu Asp Ala Ser Leu Ser Pro Val Ile Leu	
45 50 55 60	

ggc gtt caa gga gga agc cag tgc cta tct tgt ggg aca gag aaa ggg	364
Gly Val Gln Gly Gly Ser Gln Cys Leu Ser Cys Gly Thr Glu Lys Gly	
65 70 75	

cca att ctg aaa ctt gag cca gtg aac atc atg gag ctc tac ctc ggg	412
Pro Ile Leu Lys Leu Glu Pro Val Asn Ile Met Glu Leu Tyr Leu Gly	

80	85	90	
gcc aag gaa tca aag agc ttc acc ttc tac cgg cgg gat atg ggt ctt			460
Ala Lys Glu Ser Lys Ser Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu			
95	100	105	
acc tcc agc ttc gaa tcc gct gcc tac cca ggc tgg ttc ctc tgc acc			508
Thr Ser Ser Phe Glu Ser Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr			
110	115	120	
tca ccg gaa gct gac cag cct gtc agg ctc act cag atc cct gag gac			556
Ser Pro Glu Ala Asp Gln Pro Val Arg Leu Thr Gln Ile Pro Glu Asp			
125	130	135	140
ccc gcc tgg gat gct ccc atc aca gac ttc tac ttt cag cag tgt gac			604
Pro Ala Trp Asp Ala Pro Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp			
145	150	155	
tagggctgcg tgggtcccca aactccataa gcagaggcag agtaggcagt ggcggctcct			664
gatagaggat agagagacag aggagctcca cagtaggtgg cttactcttc tccttcctta			724
ctggactccc gcttctgacc taaggcacac agacactctc ttctcctgca tcccagtgtc			784
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gctggctggg ccctttccct caacctttct gacatctgca gcctctctca ttcttgccct			1084
cattctctgg ccctgaaccg agaggggtgat atcaggatag ctgacagaag atgaccaggc			1144
acactgtcct gggttgaaac cagaggggac aataaaaaac cctgattctg gtctctactc			1204
acataaaaag aagcttgtga acattaagtg ggaagagatt gctactaaat aacatacctt			1264
ggaatttcat cttaattaaa atatacttct ctatattata tatttttaaaa aaaaaaaaaa			1324
aaaaaaaaaa aaaaaaaaaa aaaaaacatg cggccg			1360

<210> 2

<211> 156

<212> PRT

<213> Mus musculus

<400> 2

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His Ala Glu Lys Val Ile Lys Gly Glu Glu Ile Ser Val Val Pro Asn	
35 40 45	
Arg Ala Leu Asp Ala Ser Leu Ser Pro Val Ile Leu Gly Val Gln Gly	
50 55 60	
Gly Ser Gln Cys Leu Ser Cys Gly Thr Glu Lys Gly Pro Ile Leu Lys	
65 70 75 80	
Leu Glu Pro Val Asn Ile Met Glu Leu Tyr Leu Gly Ala Lys Glu Ser	
85 90 95	
Lys Ser Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu Thr Ser Ser Phe	
100 105 110	
Glu Ser Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr Ser Pro Glu Ala	
115 120 125	
Asp Gln Pro Val Arg Leu Thr Gln Ile Pro Glu Asp Pro Ala Trp Asp	
130 135 140	
Ala Pro Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp	

155

<213> Mus musculus

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gaggagatca	gtgttgctcc	aaatcgggca	ctggatgcca	gtctgtcccc	tgtcatcctg	180
ggcgttcaag	gaggaagcca	gtgcctatct	tgtgggacag	agaaagggcc	aattctgaaa	240
cttgagccag	tgaacatcat	ggagctctac	ctcggggcca	aggaatcaaa	gagcttcacc	300
ttctaccggc	gggatatggg	tcttacctcc	agcttcgaat	ccgctgccta	cccaggctgg	360
ttcctctgca	cctcaccgga	agctgaccag	cctgtcaggc	tactcagat	ccctgaggac	420
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<213> Homo sapiens

<222> (57) ... (521)

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Val Leu Ser Gly Ala Leu Cys Phe Arg Met Lys Asp Ser Ala Leu Lys
5 10 15

gtg ctt tat ctg cat aat aac cag ctt cta gct gga ggg ctg cat gca 155
Val Leu Tyr Leu His Asn Asn Gln Leu Leu Ala Gly Gly Leu His Ala
20 25 30

ggg aag gtc att aaa ggt gaa gag atc agc gtg gtc ccc aat cgg tgg 203
Gly Lys Val Ile Lys Gly Glu Glu Ile Ser Val Val Pro Asn Arg Trp
35 40 45

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ctg gat gcc agc ctg tcc ccc gtc atc ctg ggt gtc cag ggt gga agc      251
Leu Asp Ala Ser Leu Ser Pro Val Ile Leu Gly Val Gln Gly Gly Ser
  50                      55                      60                      65

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cag tgc ctg tca tgt ggg gtg ggg cag gag ccg act cta aca cta gag 299
Gln Cys Leu Ser Cys Gly Val Gly Gln Glu Pro Thr Leu Thr Leu Glu
70 75 80

cca gtg aac atc atg gag ctc tat ctt ggt gcc aag gaa tcc aag agc 347
Pro Val Asn Ile Met Glu Leu Tyr Leu Gly Ala Lys Glu Ser Lys Ser
85 90 95

ttc acc ttc tac cgg cgg gac atg ggg ctc acc tcc agc ttc gag tcg 395
Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu Thr Ser Ser Phe Glu Ser

100	105	110	
gct gcc tac ccg ggc tgg ttc ctg tgc acg gtg cct gaa gcc gat cag			443
Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr Val Pro Glu Ala Asp Gln			
115	120	125	
cct gtc aga ctc acc cag ctt ccc gag aat ggt ggc tgg aat gcc ccc			491
Pro Val Arg Leu Thr Gln Leu Pro Glu Asn Gly Gly Trp Asn Ala Pro			
130	135	140	145
atc aca gac ttc tac ttc cag cag tgt gac tagggcaacg tgccccccag			541
Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp			
150	155		
aactccctgg gcagagccag ctcggtgag ggggtgagtgg aggagacca tggcggacaa			601
tcactctctc tgctctcagg acccccacgt ctgacttagt gggcacctga ccactttgtc			661
ttctggttcc cagtttggat aaattctgag atttggagct cagtccacgg tcctccccc			721
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catagagtca gggatctatg gcccttggcc cagccccacc cccttccctt taatcctgcc			1021
actgtcatat gctacctttc ctatctcttc cctcatcatc ttgttgtggg catgaggagg			1081
tggtgatgtc agaagaaatg gctcgagctc agaagataaa agataagtag ggtatgctga			1141
tcctctttta aaaacccaag atacaatcaa aatcccagat gctggtctct attcccatga			1201
aaaagtgtc atgacatatt gagaagacct acttacaag tggcatatat tgcaatttat			1261
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gc			1323

<210> 5
 <211> 155
 <212> PRT
 <213> Homo sapiens

<400> 5
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Lys Val Leu Tyr Leu His Asn Asn Gln Leu Leu Ala Gly Gly Leu His
20 25 30
Ala Gly Lys Val Ile Lys Gly Glu Ile Ser Val Val Pro Asn Arg
35 40 45
Trp Leu Asp Ala Ser Leu Ser Pro Val Ile Leu Gly Val Gln Gly Gly
50 55 60
Ser Gln Cys Leu Ser Cys Gly Val Gly Gln Glu Pro Thr Leu Thr Leu
65 70 75 80
Glu Pro Val Asn Ile Met Glu Leu Tyr Leu Gly Ala Lys Glu Ser Lys
85 90 95
Ser Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu Thr Ser Ser Phe Glu
100 105 110
Ser Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr Val Pro Glu Ala Asp
115 120 125
Gln Pro Val Arg Leu Thr Gln Leu Pro Glu Asn Gly Gly Trp Asn Ala
130 135 140
Pro Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp
145 150 155

<210> 6

<211> 465
 <212> DNA
 <213> Homo sapiens

<400> 6

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gagatcagcg	tggtcccca	tcggtggctg	gatgccagcc	tgtcccccg	catcctgggt	180
gtccaggggtg	gaagccagt	cctgtcatgt	gggggtggggc	aggagccgac	tctaactacta	240
gagccagtga	acatcatgga	gctctatctt	ggtgccaagg	aatccaagag	cttcaccttc	300
taccggcggg	acatggggct	cacctccagc	ttcgagtcgg	ctgcctaccc	gggctgggtc	360
ctgtgcacgg	tgctgaagc	cgatcagcct	gtcagactca	cccagcttcc	cgagaatggt	420
ggctggaatg	ccccatcac	agacttctac	ttccagcagt	gtgac		465

<210> 7
 <211> 177
 <212> PRT
 <213> Homo sapiens

<400> 7

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			20					25				30			
Ser	Lys	Met	Gln	Ala	Phe	Arg	Ile	Trp	Asp	Val	Asn	Gln	Lys	Thr	Phe
		35					40				45				
Tyr	Leu	Arg	Asn	Asn	Gln	Leu	Val	Ala	Gly	Tyr	Leu	Gln	Gly	Pro	Asn
	50				55						60				
Val	Asn	Leu	Glu	Glu	Lys	Ile	Asp	Val	Val	Pro	Ile	Glu	Pro	His	Ala
65					70					75					80
Leu	Phe	Leu	Gly	Ile	His	Gly	Gly	Lys	Met	Cys	Leu	Ser	Cys	Val	Lys
			85					90						95	
Ser	Gly	Asp	Glu	Thr	Arg	Leu	Gln	Leu	Glu	Ala	Val	Asn	Ile	Thr	Asp
			100					105					110		
Leu	Ser	Glu	Asn	Arg	Lys	Gln	Asp	Lys	Arg	Phe	Ala	Phe	Ile	Arg	Ser
			115				120					125			
Asp	Ser	Gly	Pro	Thr	Thr	Ser	Phe	Glu	Ser	Ala	Ala	Cys	Pro	Gly	Trp
			130				135					140			
Phe	Leu	Cys	Thr	Ala	Met	Glu	Ala	Asp	Gln	Pro	Val	Ser	Leu	Thr	Asn
145					150					155					160
Met	Pro	Asp	Glu	Gly	Val	Met	Val	Thr	Lys	Phe	Tyr	Phe	Gln	Glu	Asp
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Glu															

<210> 8
 <211> 178
 <212> PRT
 <213> Mus musculus

<400> 8

Met	Glu	Ile	Cys	Trp	Gly	Pro	Tyr	Ser	His	Leu	Ile	Ser	Leu	Leu	Leu
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			20					25					30		
Pro	Cys	Lys	Met	Gln	Ala	Phe	Arg	Ile	Trp	Asp	Thr	Asn	Gln	Lys	Thr
		35					40					45			

Phe Tyr Leu Arg Asn Asn Gln Leu Ile Ala Gly Tyr Leu Gln Gly Pro
 50 55 60
 Asn Ile Lys Leu Glu Glu Lys Ile Asp Met Val Pro Ile Asp Leu His
 65 70 75 80
 Ser Val Phe Leu Gly Ile His Gly Gly Lys Leu Cys Leu Ser Cys Ala
 85 90 95
 Lys Ser Gly Asp Asp Ile Lys Leu Gln Leu Glu Glu Val Asn Ile Thr
 100 105 110
 Asp Leu Ser Lys Asn Lys Glu Glu Asp Lys Arg Phe Thr Phe Ile Arg
 115 120 125
 Ser Glu Lys Gly Pro Thr Thr Ser Phe Glu Ser Ala Ala Cys Pro Gly
 130 135 140
 Trp Phe Leu Cys Thr Thr Leu Glu Ala Asp Arg Pro Val Ser Leu Thr
 145 150 155 160
 Asn Thr Pro Glu Glu Pro Leu Ile Val Thr Lys Phe Tyr Phe Gln Glu
 165 170 175
 Asp Gln

<210> 9
 <211> 13
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetically generated primer

<400> 9
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13

<210> 10
 <211> 17
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetically generated primer

<400> 10
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17

<210> 11
 <211> 16
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetically generated primer

<400> 11
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16

<210> 12
 <211> 16
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Synthetically generated primer

<400> 12

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16

<210> 13

<211> 2490

<212> DNA

<213> Homo sapiens

<400> 13

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<210> 14

<211> 155

<212> PRT

<213> Homo sapiens

<400> 14

Met	Val	Leu	Ser	Gly	Ala	Leu	Cys	Phe	Arg	Met	Lys	Asp	Ser	Ala	Leu
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Lys	Val	Leu	Tyr	Leu	His	Asn	Asn	Gln	Leu	Leu	Ala	Gly	Gly	Leu	His
		20						25					30		
Ala	Gly	Lys	Val	Ile	Lys	Gly	Glu	Glu	Ile	Ser	Val	Val	Pro	Asn	Arg
		35					40					45			
Trp	Leu	Asp	Ala	Ser	Leu	Ser	Pro	Val	Ile	Leu	Gly	Val	Gln	Gly	Gly
	50					55					60				
Ser	Gln	Cys	Leu	Ser	Cys	Gly	Val	Gly	Gln	Glu	Pro	Thr	Leu	Thr	Leu
65					70				75						80
Glu	Pro	Val	Asn	Ile	Met	Glu	Leu	Tyr	Leu	Gly	Ala	Lys	Glu	Ser	Lys
			85						90					95	
Ser	Phe	Thr	Phe	Tyr	Arg	Arg	Asp	Met	Gly	Leu	Thr	Ser	Ser	Phe	Glu
			100					105						110	
Ser	Ala	Ala	Tyr	Pro	Gly	Trp	Phe	Leu	Cys	Thr	Val	Pro	Glu	Ala	Asp
		115					120					125			
Gln	Pro	Val	Arg	Leu	Thr	Gln	Leu	Pro	Glu	Asn	Gly	Gly	Trp	Asn	Ala
	130					135					140				
Pro	Ile	Thr	Asp	Phe	Tyr	Phe	Gln	Gln	Cys	Asp					
145					150					155					